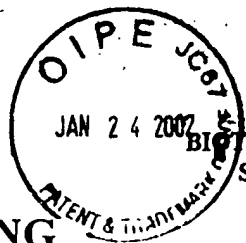
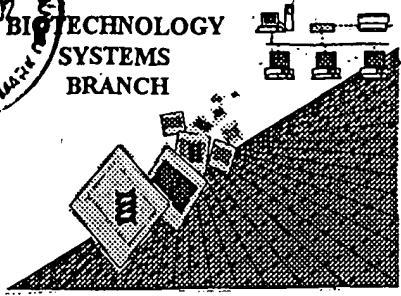


0510
1217



BIO TECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/772,134A
Source: O I P E
Date Processed by STIC: 12/18/01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE: SEE BELOW FOR ADDRESS:
<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by the treatment given to all mail coming via the Brentwood Mail Facility.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom, including:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>) , EFS Submission User Manual - ePAVE)
2. U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
1911 South Clark Street, Crystal Mall One, Sequence Information, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, 2011 South Clark Place, Customer Window, Box Sequence, Crystal Plaza Two,
Lobby, Room 1B03, Arlington, Virginia 22202
4. Federal Express Delivery, 2011 South Clark Street, Crystal Plaza 2, Room 1B03-Mailroom, Box Sequence,
Arlington, VA 22202

Raw Sequence Listing Error Summary .

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 091772,134A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.

- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

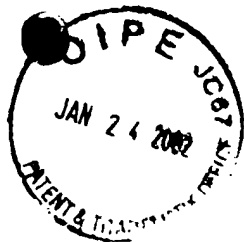
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000

- 9 ✓ Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.



O I P E

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/772,134A

DATE: 12/18/2001
TIME: 10:27:06

Input Set : A:\seq list 1268-4-2.ST25.txt
Output Set: N:\CRF3\12182001\I772134A.raw

**Does Not Comply
Corrected Diskette Needed**

Errors on pp. 3+4

3 <110> APPLICANT: Lightfoot, David
4 Meksem, Khalid
6 <120> TITLE OF INVENTION: ISOLATED POLYNUCLEOTIDES AND POLYPEPTIDES RELATING TO LOCI
7 UNDERLYING RESISTANCE TO SOYBEAN CYST NEMATODE AND SOYBEAN
8 SUDDEN DEATH SYNDROME AND METHODS EMPLOYING SAME
10 <130> FILE REFERENCE: 1268/4/2
:--> 12 <140> CURRENT APPLICATION NUMBER: US/09/772,134A
:--> 12 <141> CURRENT FILING DATE: 2001-12-18
12 <150> PRIOR APPLICATION NUMBER: 60/178,811
13 <151> PRIOR FILING DATE: 2000-01-28
15 <160> NUMBER OF SEQ ID NOS: 122
17 <170> SOFTWARE: PatentIn version 3.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 87
21 <212> TYPE: DNA
22 <213> ORGANISM: soybean
24 <400> SEQUENCE: 1
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27 tgggtttctg ataaccgtgg tcgttaa 87
30 <210> SEQ ID NO: 2
31 <211> LENGTH: 92
32 <212> TYPE: DNA
33 <213> ORGANISM: soybean
35 <400> SEQUENCE: 2
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38 agatttgggt ttcagataac cgtggctcgt aa 92
41 <210> SEQ ID NO: 3
42 <211> LENGTH: 113
43 <212> TYPE: DNA
44 <213> ORGANISM: soybean
46 <400> SEQUENCE: 3
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52 <210> SEQ ID NO: 4
53 <211> LENGTH: 135
54 <212> TYPE: DNA
55 <213> ORGANISM: soybean
57 <400> SEQUENCE: 4
58 ttaaagggat atgttttttt cactaatgct gtaaaaattc acccagattt ttgcattttc 60
60 tttgaaaaaa tgtactagat atatcatgtt tttttacaag cattacaata atattcactc 120
62 gtatattagg aattc 135
65 <210> SEQ ID NO: 5
66 <211> LENGTH: 116
67 <212> TYPE: DNA
68 <213> ORGANISM: soybean
70 <400> SEQUENCE: 5
71 gaattccggt tatctcagac aacttttgggt tggtttgggt atagtaaaga cacgattatc 60

end



The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

RAW SEQUENCE LISTING

DATE: 12/18/2001

PATENT APPLICATION: US/09/772,134A

TIME: 10:27:06

Input Set : A:\seq list 1268-4-2.ST25.txt

Output Set: N:\CRF3\12182001\I772134A.raw

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77 <211> LENGTH: 113
78 <212> TYPE: DNA
79 <213> ORGANISM: soybean
81 <400> SEQUENCE: 6
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84 caggctttga gaggcataga aataattttt ttatataaaa aaaagtctct tta      113
87 <210> SEQ ID NO: 7
88 <211> LENGTH: 409
89 <212> TYPE: DNA
90 <213> ORGANISM: soybean - Forrest
92 <400> SEQUENCE: 7
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97 aactcaggta tcaatctaaa agcaacgctg ttcaccttga gctgaaacac ctggaggaga      180
99 aagcaaagca aaccaaagcg gagagagaaa taaagaacgg aaacagagag agagagagga      240
101 aggaccttgt tcaaagcaac ggggacaact ttagagccct ggcgcgcgtg ggggtcaata      300
103 agcgtaacct ggctgaggag agcctcggcg tcgtccttgc tgaagcagaa gaggaagagc      360
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109 <211> LENGTH: 417
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111 <213> ORGANISM: soybean
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118 ttgacttgag agaactcagg tatcaatcta aaagcaacgc tgttcacctt gagctgaaac      180
120 acctggagga gaaagcaaag caaaccaaac gcgagagaga aataaagaac ggaaacagag      240
122 agagaggaag gacctgttc aaagcaacgg ggacaacttt agagccctgg cgcgcgtggg      300
124 ggtcaataag cgtaacctgg ctgaggagag cctcggcgcc gtccttgctg aagcagaaga      360
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131 <212> TYPE: DNA
132 <213> ORGANISM: soybean
134 <400> SEQUENCE: 9
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143 <211> LENGTH: 164
144 <212> TYPE: DNA
145 <213> ORGANISM: soybean
147 <400> SEQUENCE: 10
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150 ctttttttta ttgaaaaaat cgacccaagt tgaaacacat gtttgagaat tgttttggtg      120
152 atccaacgtt tttcttgta aatcagctgt gagaggggaa ttgg      164
155 <210> SEQ ID NO: 11

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RAW SEQUENCE LISTING

DATE: 12/18/2001

PATENT APPLICATION: US/09/772,134A

TIME: 10:27:06

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Output Set: N:\CRF3\12182001\I772134A.raw

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 158 <213> ORGANISM: soybean
 160 <400> SEQUENCE: 11

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163 ggaactgcta ttcttacttc taaatttttt attgacatcc aaagtgtgct ttaa	114

166 <210> SEQ ID NO: 12
 167 <211> LENGTH: 114
 168 <212> TYPE: DNA
 169 <213> ORGANISM: soybean
 171 <400> SEQUENCE: 12

172 gaattcccag ccagatttgt atcaaacatg tattgtccac aaaatgttca agcatcttag	60
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177 <210> SEQ ID NO: 13
 178 <211> LENGTH: 3106
 179 <212> TYPE: DNA
 180 <213> ORGANISM: soybean
 182 <220> FEATURE:
 183 <221> NAME/KEY: misc_feature
 184 <223> OTHER INFORMATION: n is an undetermined nucleotide (dATP, dCTP, dGTP, or dTTP)
 187 <400> SEQUENCE: 13

188 aatgggagga gtgggaaaga cagtggctat ggagcttggt cggagggttg ggttggaatc	60
190 aagtgtgctc agggacaggt tattgtgatc cagcttcctt ggaagggttt gaggggtcga	120
192 atcaccgaca aaattggcca acttcaaggc ctcaggaagc ttagtcttca tgataaccaa	180
194 attggtgggt caatcccttc aactttggga cttcttccca accttagagg ggttcagtta	240
196 ttcaacaata ggcttacagg ttccatacct ctttcttttag gtttctgcct ttgcttcaag	300
198 tctcttgacc tcagcaacaa cttgtctaca ggagcaatcc cttatagtct tgctaattcc	360
200 actaagcttt attggcttaa cttgagtttc aaactcctct ctggctcctt accagctagc	420
202 ctaactcact cattttctct cacttttctt tctcttcaaa ataacaatct ttctggctcc	480
204 cttcctaact cttgggggtg gaattccaag aatggcttct ttaggcttca aaatttgatc	540
206 ctatgcata actttttcac tgggtgacgt cctgcttctt tgggtagctt aagagagctc	600
208 aatgagattt cccttagtca taataagttt agtggagcta taccaaatga aataggaacc	660
210 ctttctaggc ttaagacact tgacatttct aataatgcct tgaatgggaa cttgcctgct	720
212 accctctcta atttatectc acttacactg ctgaatgcag agaacaacct ccttgacaat	780
214 caaatccctc aaagtttagg tagattgcgt aatctttctg ttctgatttt gagtagaaac	840
216 caatttagtg gacatattcc ttcaagcatt gcaaacattt cctcgcttag gcagcttgat	900
218 ttgtcactga ataatttcag tggagaaatt ccagttctct ttgacagtca gcgcagtcta	960
220 aatctcttca atgtttccta caatagcctc tcaggttctg tccccctct gcttgccaag	1020
222 aaatttaact caagctcatt tgtgggaaat attcaactat gtgggtacag cccttcaacc	1080
224 ccatgtcttt cccaagctcc atcacaagga gtcattgcc cacctcctga agtgtcaaaa	1140
226 catcaccatc ataggaagct aagcaccaaa gacataattc tcatagtagc aggagttctc	1200
228 ctctagatcc tgattatact ttgttgtgtc ctgcttttct gcctgatcag aaagagatca	1260
230 acatctaggc cgggaacggc caagccaccc gagggtagag cggccactat gaggacagaa	1320
232 aaaggagtcc ctccagttgc tgggtggtgat gttgaagcag gtggggaggc tggagggaaa	1380
234 ctagtccatt ttgatggacc aatggctttt acagctgatg atctcttggt tgoaacagct	1440
236 gagatcatgg gaaagagcac ctatggaact gtttataagg ctattttgga ggatggaaagt	1500
238 caagttgcag taaagagatt gagggaaaag atcactaaaag gtcatagaga atttgaatca	1560
240 gaagtcagtg ttctaggaaa aattagacac cccaatgttt tggtctgag ggcctattac	1620
242 ttgggacca aaggggaaaa gcttctgggt ttgtatacat gtctaaagga agtcttgctt	1680

RAW SEQUENCE LISTING

DATE: 12/18/2001

PATENT APPLICATION: US/09/772,134A

TIME: 10:27:06

Input Set : A:\seq list 1268-4-2.ST25.txt

Output Set: N:\CRF3\12182001\I772134A.raw

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244 ctttcctaca tggaagggtc gtgtgctggt tctttcatta aagtgttggt tgtgctgggc 1740
246 tttaattata atttgaggtt ttaccttagt aatctgtata attctaactg gagaacagta 1800
v--> 248 caaacaaaaa cacctaagga acaacacctt anctttaata taccatatca ataaagttaa 1860
250 atattttctt ggtcatcttg atgcaggggg aactgaacat tcattattgg ccacaagatt 1920
252 aaaatagccc aagccttggc cggggcttgt ttgccttcac tcccaggaga acatcataca 1980
v--> 254 tgggacctcn catccagcaa tgtgtggcct gatgaaaaac aaatgctaaa attcagattt 2040
256 tggctctttt cgggttgatg tcaactgctg ctaattccaa cgtgatagct acagctggag 2100
258 cattggatac cgggcacctg agctctcaaa gctcaagaaa gcaaacta aaactgatat 2160
260 ctacagtctt ggtgttatct tgttagaact cctaacgagg aaatcacctg ggggtgtctat 2220
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264 ggtttttgat gcagacttga tgagagatgc atccacagtt ggcgacgagt tgctaaacac 2340
266 gttgaagctc gctttgcact gtgttgatcc ttctccatca gcacgaccag aagttcatca 2400
268 agttctccag cagctgaaga gattagacca gagagatcag tcacagccag tcccggggac 2460
270 gatatcgtat agcacaatt ttgcattgat tttttgtgc caaatgtagt aggcctacta 2520
272 tatatatgtt ctatgattct ttcattctta tattattttt gcctgtttga atgcttgaat 2580
274 ttgtacatac tcatactaca ataagggtga gttctggtta attttacctc tacctcaaag 2640
276 ctgggggtga attctgtttc ctccaaggca cataatagtt gaaaatagtt ctcaggagca 2700
278 ttcattgttt attctgcaag attctctttc acggtgcta tcttctatgc atgccctgcc 2760
280 cataaatgca ttatgaagaa ttgtaacggc tgtgtttttg gacttcttca aaaagtttat 2820
282 gttattgcca ggtgtatata tcaacatgtt ttaaagattt tcaaacaatc aggttttaga 2880
284 tgtgggtttg catgcatgag attggactag tgcgcttgat gtagtataaa atataaattg 2940
286 tccaatcaag caccctctac atgtccaaat aatgggcctt atgaaactta attttttaat 3000
i--> 288 tacaactac agtaatcttt ttgaataaag atttacaat tacaacngac atgtgaagen 3060
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293 <210> SEQ ID NO: 14
294 <211> LENGTH: 830
295 <212> TYPE: PRT
296 <213> ORGANISM: soybean
298 <220> FEATURE:
299 <221> NAME/KEY: misc_feature
300 <223> OTHER INFORMATION: X is any amino acid
303 <400> SEQUENCE: 14
305 Asn Gly Arg Ser Gly Lys Asp Ser Gly Tyr Gly Ala Cys Ser Gly Gly
306 1 5 10 15
308 Trp Val Gly Ile Lys Cys Ala Gln Gly Gln Val Ile Val Ile Gln Leu
309 20 25 30
311 Pro Trp Lys Gly Leu Arg Gly Arg Ile Thr Asp Lys Ile Gly Gln Leu
312 35 40 45
314 Gln Gly Leu Arg Lys Leu Ser Leu His Asp Asn Gln Ile Gly Gly Ser
315 50 55 60
317 Ile Pro Ser Thr Leu Gly Leu Leu Pro Asn Leu Arg Gly Val Gln Leu
318 65 70 75 80
320 Phe Asn Asn Arg Leu Thr Gly Ser Ile Pro Leu Ser Leu Gly Phe Cys
321 85 90 95
323 Pro Leu Leu Gln Ser Leu Asp Leu Ser Asn Asn Leu Leu Thr Gly Ala
324 100 105 110
326 Ile Pro Tyr Ser Leu Ala Asn Ser Thr Lys Leu Tyr Trp Leu Asn Leu
327 115 120 125
329 Ser Phe Asn Ser Phe Ser Gly Pro Leu Pro Ala Ser Leu Thr His Ser

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*must give location of X22
in 2227, see error
summary sheet, Term 9*

RAW SEQUENCE LISTING

DATE: 12/18/2001

PATENT APPLICATION: US/09/772,134A

TIME: 10:27:06

Input Set : A:\seq list 1268-4-2.ST25.txt

Output Set: N:\CRF3\12182001\I772134A.raw

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332 Phe Ser Leu Thr Phe Leu Ser Leu Gln Asn Asn Asn Leu Ser Gly Ser
333 145      150      155      160
335 Leu Pro Asn Ser Trp Gly Gly Asn Ser Lys Asn Gly Phe Phe Arg Leu
336      165      170      175
338 Gln Asn Leu Ile Leu Asp His Asn Phe Thr Gly Asp Val Pro Ala
339      180      185      190
341 Ser Leu Gly Ser Leu Arg Glu Leu Asn Glu Ile Ser Leu Ser His Asn
342      195      200      205
344 Lys Phe Ser Gly Ala Ile Pro Asn Glu Ile Gly Thr Leu Ser Arg Leu
345      210      215      220
347 Lys Thr Leu Asp Ile Ser Asn Asn Ala Leu Asn Gly Asn Leu Pro Ala
348 225      230      235      240
350 Thr Leu Ser Asn Leu Ser Ser Leu Thr Leu Leu Asn Ala Glu Asn Asn
351      245      250      255
353 Leu Leu Asp Asn Gln Ile Pro Gln Ser Leu Gly Arg Leu Arg Asn Leu
354      260      265      270
356 Ser Val Leu Ile Leu Ser Arg Asn Gln Phe Ser Gly His Ile Pro Ser
357      275      280      285
359 Ser Ile Ala Asn Ile Ser Ser Leu Arg Gln Leu Asp Leu Ser Leu Asn
360      290      295      300
362 Asn Phe Ser Gly Glu Ile Pro Val Ser Phe Asp Ser Gln Arg Ser Leu
363 305      310      315      320
365 Asn Leu Ser Asn Val Ser Tyr Asn Ser Leu Ser Gly Ser Val Pro Pro
366      325      330      335
368 Leu Leu Ala Lys Lys Phe Asn Ser Ser Ser Phe Val Gly Asn Ile Gln
369      340      345      350
371 Leu Cys Gly Tyr Ser Pro Ser Thr Pro Cys Leu Ser Gln Ala Pro Ser
372      355      360      365
374 Gln Gly Val Ile Ala Pro Pro Glu Val Ser Lys His His His His
375      370      375      380
377 Arg Lys Leu Ser Thr Lys Asp Ile Ile Leu Ile Val Ala Gly Val Leu
378 385      390      395      400
380 Leu Val Val Leu Ile Ile Leu Cys Cys Val Leu Leu Phe Cys Leu Ile
381      405      410      415
383 Arg Lys Arg Ser Thr Ser Lys Ala Gly Asn Gly Gln Ala Thr Glu Gly
384      420      425      430
386 Arg Ala Ala Thr Met Arg Thr Glu Lys Gly Val Pro Pro Val Ala Gly
387      435      440      445
389 Gly Asp Val Glu Ala Gly Gly Glu Ala Gly Gly Lys Leu Val His Phe
390      450      455      460
392 Asp Gly Pro Met Ala Phe Thr Ala Asp Asp Leu Leu Cys Ala Thr Ala
393 465      470      475      480
395 Glu Ile Met Gly Lys Ser Thr Tyr Gly Thr Val Tyr Lys Ala Ile Leu
396      485      490      495
398 Glu Asp Gly Ser Gln Val Ala Val Lys Arg Leu Arg Glu Lys Ile Thr
399      500      505      510
401 Lys Gly His Arg Glu Phe Glu Ser Glu Val Ser Val Leu Gly Lys Ile
402      515      520      525

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VERIFICATION SUMMARY

DATE: 12/18/2001

PATENT APPLICATION: US/09/772,134A

TIME: 10:27:07

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Output Set: N:\CRF3\12182001\I772134A.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:248 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:13
L:248 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:254 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:13
L:254 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:288 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:13
L:288 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:290 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:13
L:290 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:419 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:419 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:422 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:422 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:425 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:425 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:428 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:428 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:434 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:434 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:437 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:437 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:446 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:446 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:452 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:452 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:455 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:455 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:467 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:15
L:475 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:487 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:493 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:495 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:497 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:506 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:532 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:541 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:17
L:547 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:549 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:551 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
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L:559 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:561 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:563 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:565 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:574 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:18

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/772,134A

DATE: 12/18/2001

TIME: 10:27:07

Input Set : A:\seq list 1268-4-2.ST25.txt

Output Set: N:\CRF3\12182001\I772134A.raw

.:596 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
.:598 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
.:600 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
.:602 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
.:604 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
.:615 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:19
.:633 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
.:637 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
.:639 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
.:641 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
.:643 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
.:645 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
.:683 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:20
.:683 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
.:685 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:20
.:685 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
.:687 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:20
.:687 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
.:689 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:20
.:689 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
.:717 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:21
.:717 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
.:719 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:21
.:719 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
.:721 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:21
.:721 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
.:723 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:21
.:723 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
.:725 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:21
.:725 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
.:727 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:21
.:727 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
.:755 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:22
.:759 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:22
.:761 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:22
.:763 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:22
.:765 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:22
.:767 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:22
.:769 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:22
.:799 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:23
.:803 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:23
.:805 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:23
.:807 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:23
.:809 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:23
.:837 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:24
.:839 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:24
.:841 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:24
.:877 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:25
.:879 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:25

VERIFICATION SUMMARY

DATE: 12/18/2001

PATENT APPLICATION: US/09/772,134A

TIME: 10:27:07

Input Set : A:\seq list 1268-4-2.ST25.txt

Output Set: N:\CRF3\12182001\I772134A.raw

L:893 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:26
L:909 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:26
L:911 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:26
L:945 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:27
L:961 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:28
L:963 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:28
L:975 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:28
L:977 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:28
L:979 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:28
L:981 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:28

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CProject    ~~
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CDNASequence      E-ATG/ M-CG87 Forrestsoybean"gaattcatgg
tttctcttat gacattgttg ccaagtaata ctactatata aattcagatt      60
tgggtttctg ataaccgtgg tcgttaa
87      agctrymkswbdhvn      DNA      ~~      CCom
mentFeature      ¥      ?      E-ATG/ M-CGA92
Essexsoybean\gaattcatggtttctcttatcttatgacattgttgccaagtaatactactat
ataaattcagatttgggtttcagataaccgtggctcgtaa      agctrymkswbdhvn
DNA      ?
      □      ?      E-CTA/ M-AGG113
Forrestsoybean-gaattcctaa tatacagagtg aatattattg taatgcttgt
aaaaaaacat gataaaatgc      60
aaaaatttgg ggtgaatttt tacgacatta gtgaaaaaaaa catatccctt taa
113
      agctrymkswbdhvn      DNA      ?
      OF'      ?      E-CTA/ M-AGG135
Essexsoybeanatttaaagggat atgtttttt cactaatgct gtaaaaattc
accagattt ttgcatttc      60
tttgaaaaaa tgtactagat atatcatgtt tttttacaag cattacaata atattcactc
120
gtatattagg aattc
135      agctrymkswbdhvn      DNA      ?
      gtga      ?      E-CGG/ M-AGA116 Forrestsoybean-gaattccggt
tatctcagac aacttttgtt tggtttgggt atagtaaaga caccgattatc      60

```

Met Val Ser Met Phe Ser Leu Ser Phe Lys Trp Pro Gly Phe Cys Leu

1

5

10

15

Phe Val

| Numeric Identifier | Definition                    | Comments and format                                                                                      | Mandatory (M) or optional (O)                                          |
|--------------------|-------------------------------|----------------------------------------------------------------------------------------------------------|------------------------------------------------------------------------|
| <110>              | Applicant                     | Preferably max. of 10 names; one name per line; preferable format: Surname, Other Names and/or Initials. | M                                                                      |
| <120>              | Title of Invention            |                                                                                                          | M                                                                      |
| <130>              | File Reference                | Personal file reference                                                                                  | M when filed prior to assignment of appl. number.                      |
| <140>              | Current Application Number    | Specify as: US 07/999,999 or PCT/US96/99999                                                              | M, if available.                                                       |
| <141>              | Current Filing Date           | Specify as: yyyy-mm-dd                                                                                   | M, if available.                                                       |
| <150>              | Prior Application Number      | Specify as: US 07/999,999 or PCT/US96/99999                                                              | M, if applicable. Include priority documents under 35 USC 119 and 120. |
| <151>              | Prior Application Filing Date | Specify as: yyyy-mm-dd                                                                                   | M, if applicable.                                                      |
| <160>              | Number of SEQ ID NOs          | Count includes total number of SEQ ID NOs                                                                | M.                                                                     |
| <170>              | Software                      | Name of software used to create the Sequence Listing.                                                    | O.                                                                     |
| <210>              | SEQ ID NO:1                   | Response shall be an integer representing the SEQ ID NO shown.                                           | M.                                                                     |
| <211>              | Length                        | Respond with an integer expressing the number of bases or amino acid residues.                           | M.                                                                     |

| Numeric Identifier | Definition                | Comments and format                                                                                                                                                                                                                                                    | Mandatory (M) or optional (O)                                                                                                                                                                                        |
|--------------------|---------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <212>              | Type                      | Whether presented sequence molecule is DNA, RNA, or PRT (protein). If a nucleotide sequence contains both DNA and RNA fragments, the type shall be "DNA." In addition, the combined DNA/RNA molecule shall be further described in the <220> to <223> feature section. | M.                                                                                                                                                                                                                   |
| <213>              | Organism                  | Scientific name, i.e. Genus/ species, Unknown or Artificial Sequence. In addition, the "Unknown" or "Artificial Sequence" organisms shall be further described in the <220> to <223> feature section.                                                                  | M                                                                                                                                                                                                                    |
| <220>              | Feature                   | Leave blank after <220>. <221-223> provide for a description of points of biological significance in the sequence..                                                                                                                                                    | M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified base was used in a sequence; if ORGANISM is "Artificial Sequence" or "Unknown"; if molecule is combined DNA/RNA  |
| <221>              | Name/Key                  | Provide appropriate Identifier for feature, preferably from WIPO Standard ST.25 (1998), Appendix 2, Tables 5 and 6.                                                                                                                                                    | M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified base was used in a sequence.                                                                                     |
| <222>              | Location                  | Specify location within sequence; where appropriate state number of first and last bases/ amino acids in feature.                                                                                                                                                      | M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified base was used in a sequence.                                                                                     |
| <223>              | Other Information         | Other relevant information; four lines maximum                                                                                                                                                                                                                         | M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified base was used in a sequence; if ORGANISM is "Artificial Sequence" or "Unknown"; if molecule is combined DNA/RNA. |
| <300>              | Publication Information   | Leave blank after <300>                                                                                                                                                                                                                                                | O.                                                                                                                                                                                                                   |
| <301>              | Authors                   | Preferably max of ten named authors of publication; specify one name per line; preferable format: Surname, Other Names and/or Initials.                                                                                                                                | O.                                                                                                                                                                                                                   |
| <302>              | Title                     |                                                                                                                                                                                                                                                                        | O.                                                                                                                                                                                                                   |
| <303>              | Journal                   |                                                                                                                                                                                                                                                                        | O.                                                                                                                                                                                                                   |
| <304>              | Volume                    |                                                                                                                                                                                                                                                                        | O.                                                                                                                                                                                                                   |
| <305>              | Issue                     |                                                                                                                                                                                                                                                                        | O.                                                                                                                                                                                                                   |
| <306>              | Pages                     |                                                                                                                                                                                                                                                                        | O.                                                                                                                                                                                                                   |
| <307>              | Date                      | Journal date on which data published; specify as yyyy-mm-dd, MMM-yyyy or Season-yyyy.                                                                                                                                                                                  | O.                                                                                                                                                                                                                   |
| <308>              | Database Accession Number | Accession number assigned by database including database name.                                                                                                                                                                                                         | O.                                                                                                                                                                                                                   |
| <309>              | Database Entry Date       | Date of entry in database; specify as yyyy-mm-dd or MMM-yyyy.                                                                                                                                                                                                          | O.                                                                                                                                                                                                                   |
| <310>              | Patent Document Number    | Document number; for patent-type citations only. Specify as, for example, US 07/999,999.                                                                                                                                                                               | O.                                                                                                                                                                                                                   |
| <311>              | Patent Filing Date        | Document filing date, for patent-type citations only; specify as yyyy-mm-dd.                                                                                                                                                                                           | O.                                                                                                                                                                                                                   |
| <312>              | Publication Date          | Document publication date, for patent-type citations only; specify as yyyy-mm-dd.                                                                                                                                                                                      | O.                                                                                                                                                                                                                   |
| <313>              | Relevant Residues         | FROM (position) TO (position)                                                                                                                                                                                                                                          | O.                                                                                                                                                                                                                   |
| <400>              | Sequence                  | SEQ ID NO should follow the numeric identifier and should appear on the line preceding the actual sequence.                                                                                                                                                            | M.                                                                                                                                                                                                                   |